

# Role of Genetics in Cancer Control and Public Health



INOVA TRANSLATIONAL  
MEDICINE INSTITUTE

— individualized care through science and technology

20<sup>th</sup> Annual Maryland

Council on Cancer Control Conference

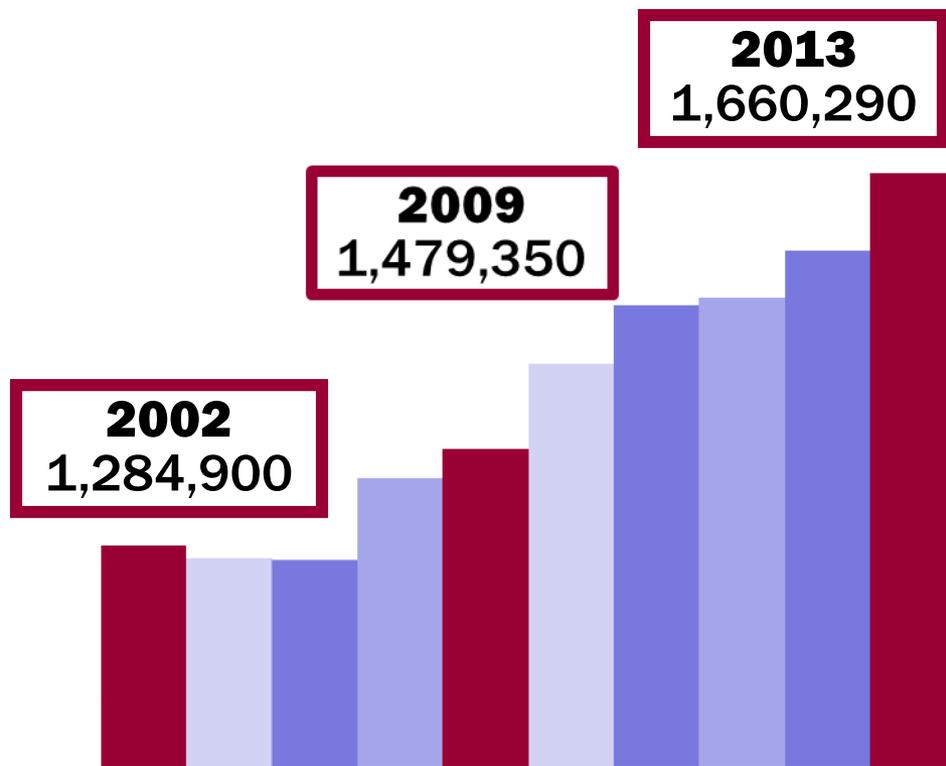
November 19, 2013



INOVA™ HEALTH  
SYSTEM

# Why What We Are Doing is So Important

## Human and Economic Burden of Cancer



Estimated # of New U.S. Cancer Cases

- **1,660,290** Americans will be diagnosed in 2013
- **580,350 are expected to die of cancer in 2013**
- More than **1,600 people a day die from cancer**
- **Cost of cancer care was \$157 billion in 2012**
- Second most common cause of death

# Cancer Cases Are Rising Globally

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25,000,000

20,000,000

15,000,000

10,000,000

5,000,000

- Today, more than half of new cancer cases and nearly two-thirds of cancer deaths occur in the low income, lower middle income, and upper middle income countries of the developing world
- **By 2030, the developing world is expected to bear 70% of the global cancer burden**

# A Transformation in Medicine

## Chronic Diseases Account for >65% of Health Care Costs.

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### Aging Population

- The fastest growing component of the U.S. population is the age group > 75 years.
- Those > 65 years are more likely to have more than one chronic disease problem.

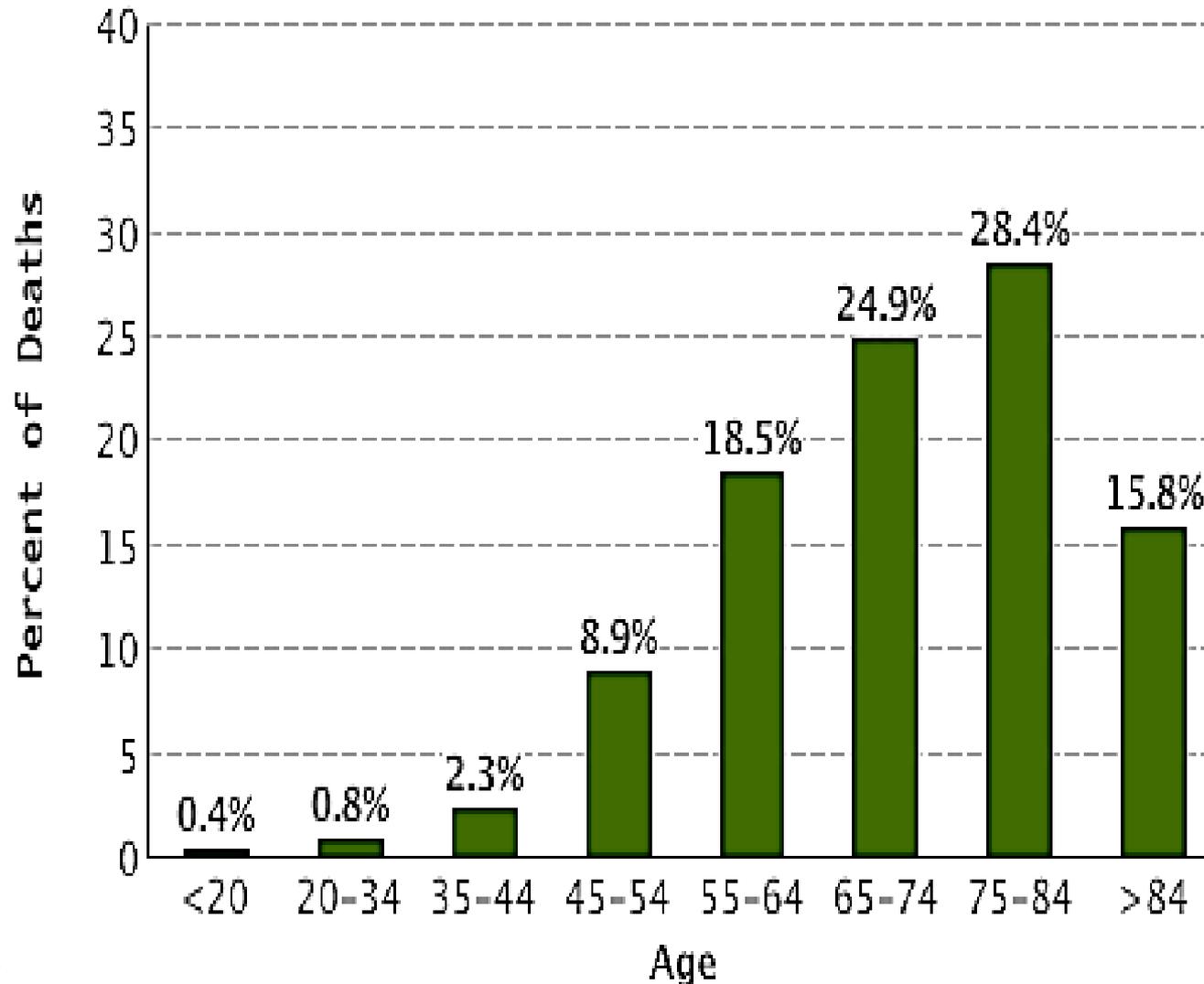
### Life Style and Behavior

- Chronic diseases often share man-made causes – tobacco, obesity, substance abuse and inactivity.

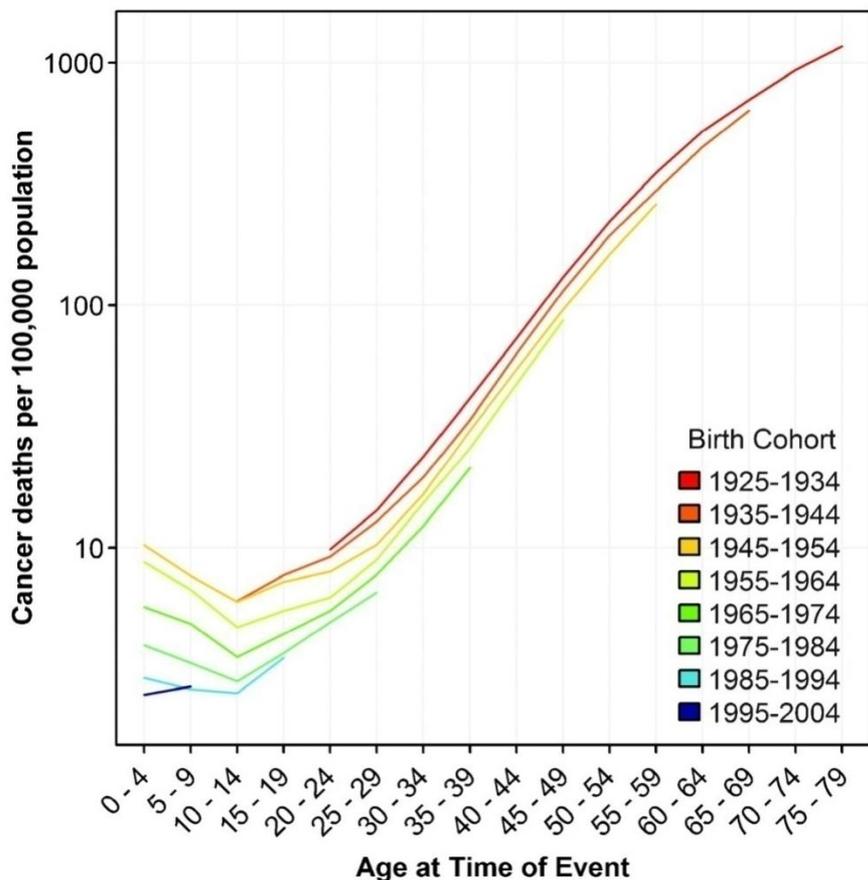
### Health Disparities

- Chronic diseases are often a greater burden in the poor and less educated population
- Access to care will be an increasing determinant of cost and of disease mortality

# All-Site Cancer Death Rates by Age

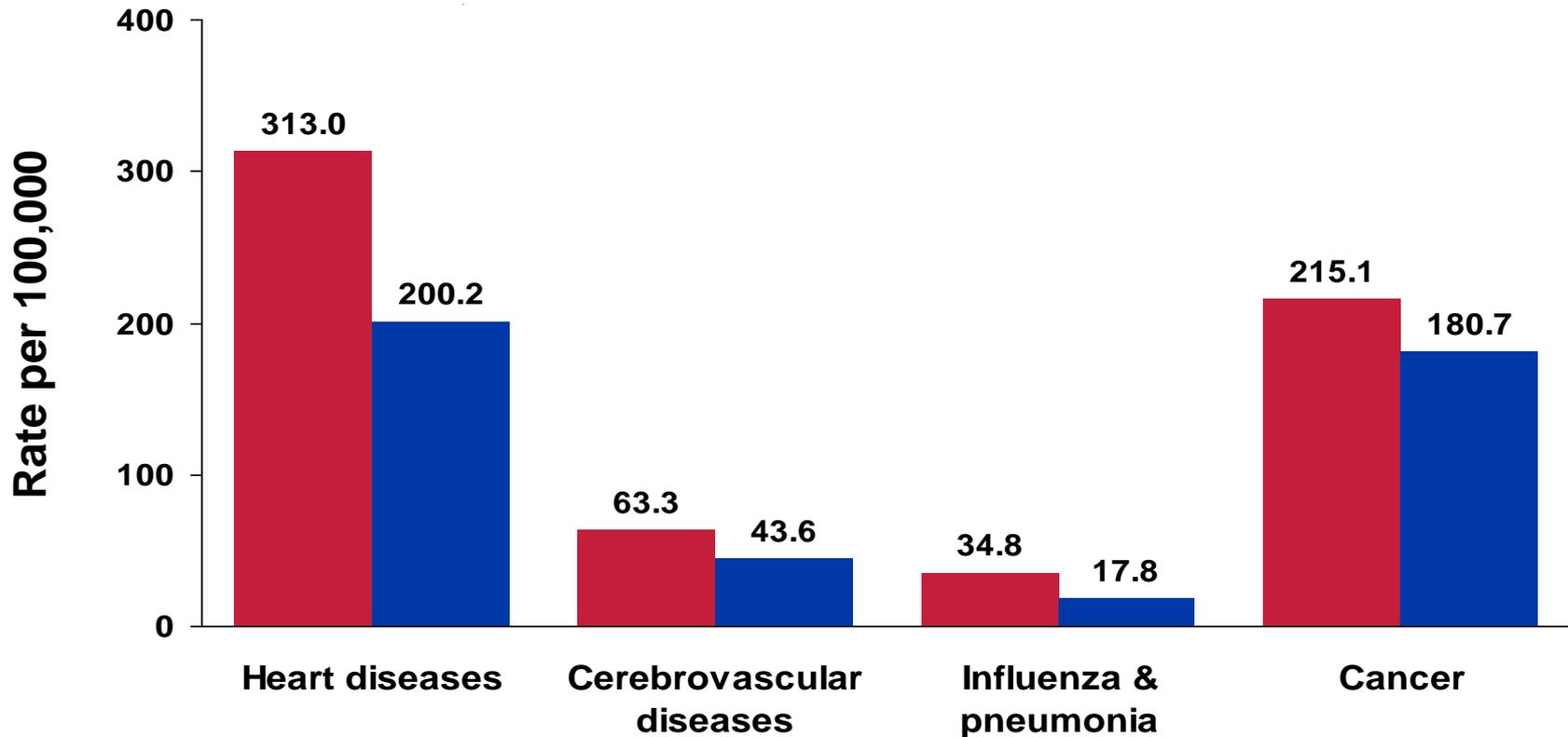


# All-Site Cancer Rates in Successive Birth Cohorts by Age of Death



- Mortality rates for cancer in the U.S. have declined over past 50 years despite relatively stable incidence (except lung cancer)
- Declines suggest that better cancer detection, treatment and prevention have been effective
- **Effects may be even larger than currently observed**; cancer death rates at every age have been successively lower for each generation since 1925
- As younger groups age, their lower death rates will greatly impact (reduce) age-adjusted death rates utilized to mark our progress against cancer

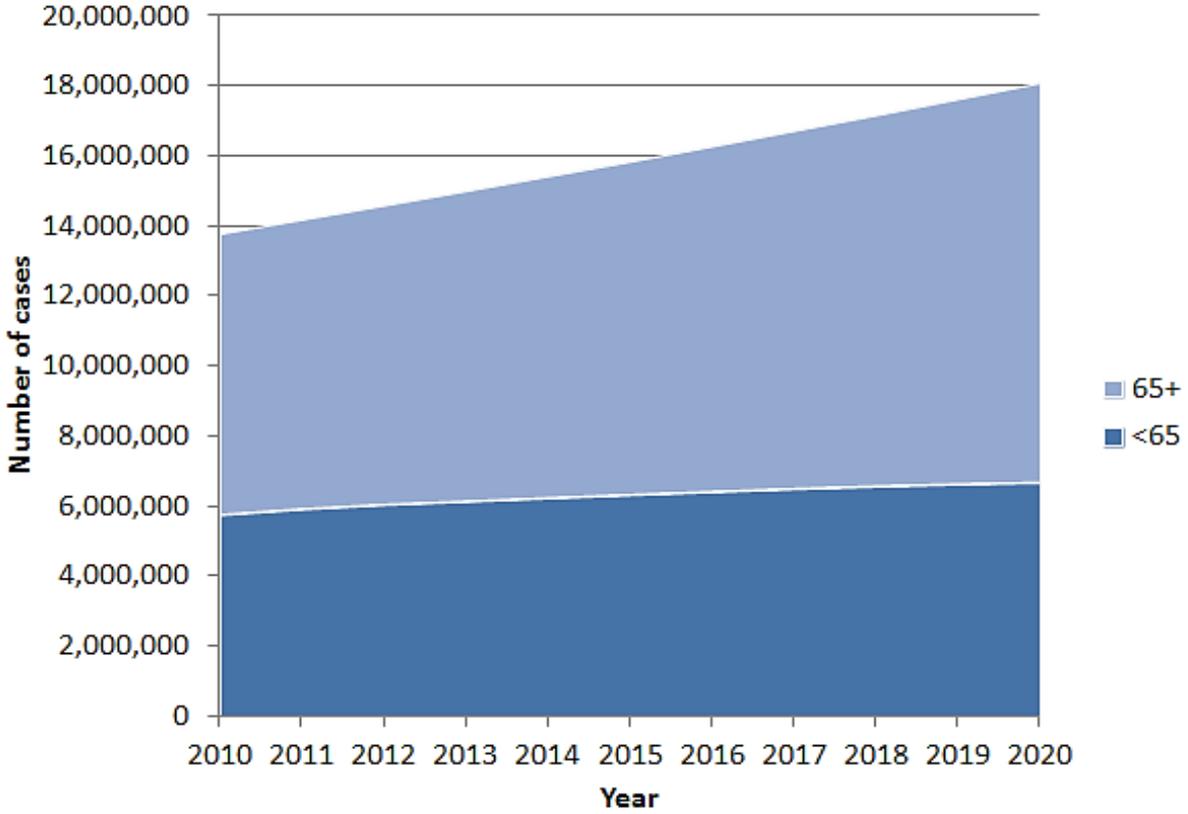
# Declining U.S. Mortality Rates from 1991-2006



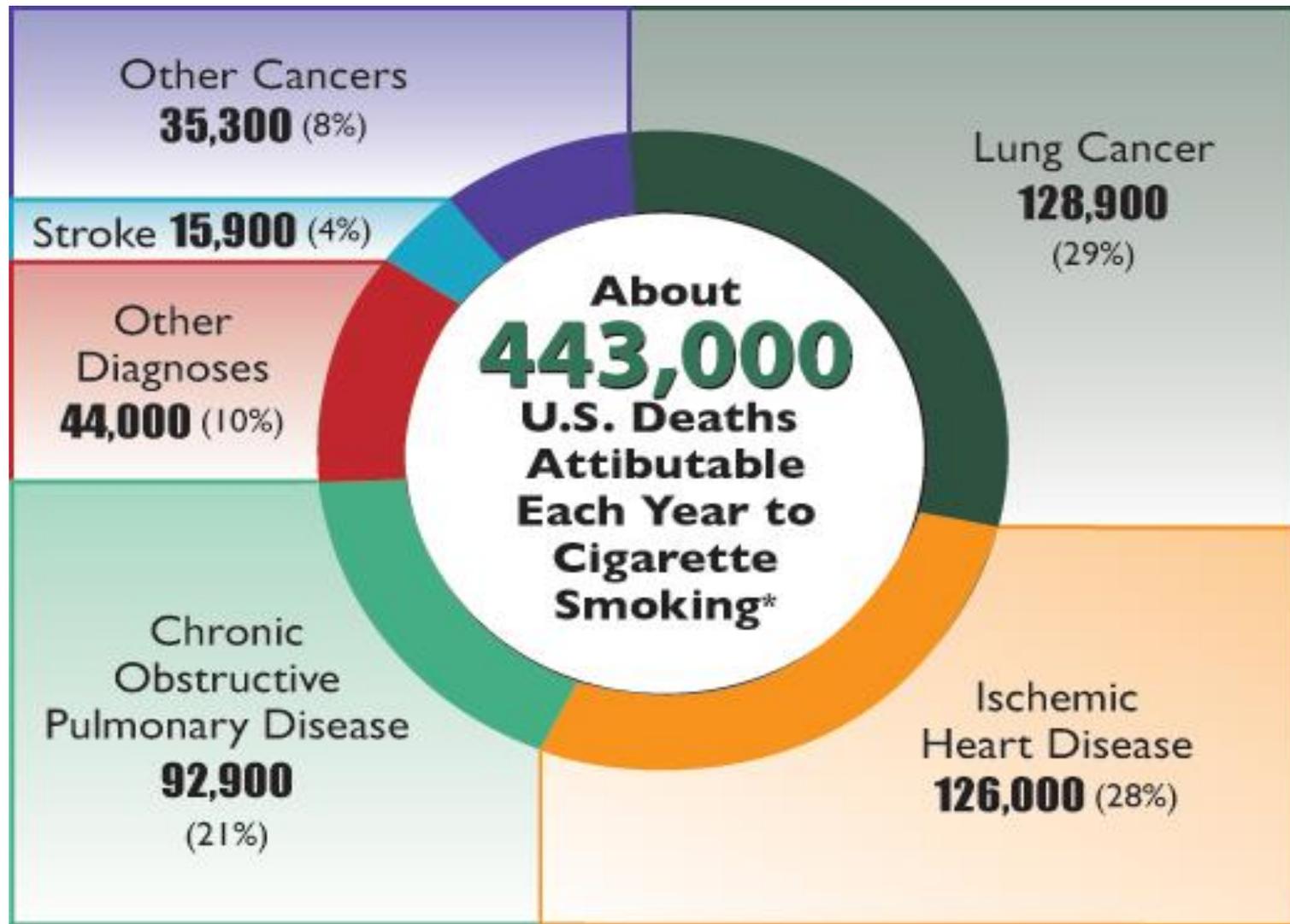
Rates are age-adjusted to 2000 U.S. standard population

U.S. mortality data, National Center for Health Statistics, Centers  
for Disease Control and Prevention, 2009

# Growth in the Population of Cancer Survivors by Age

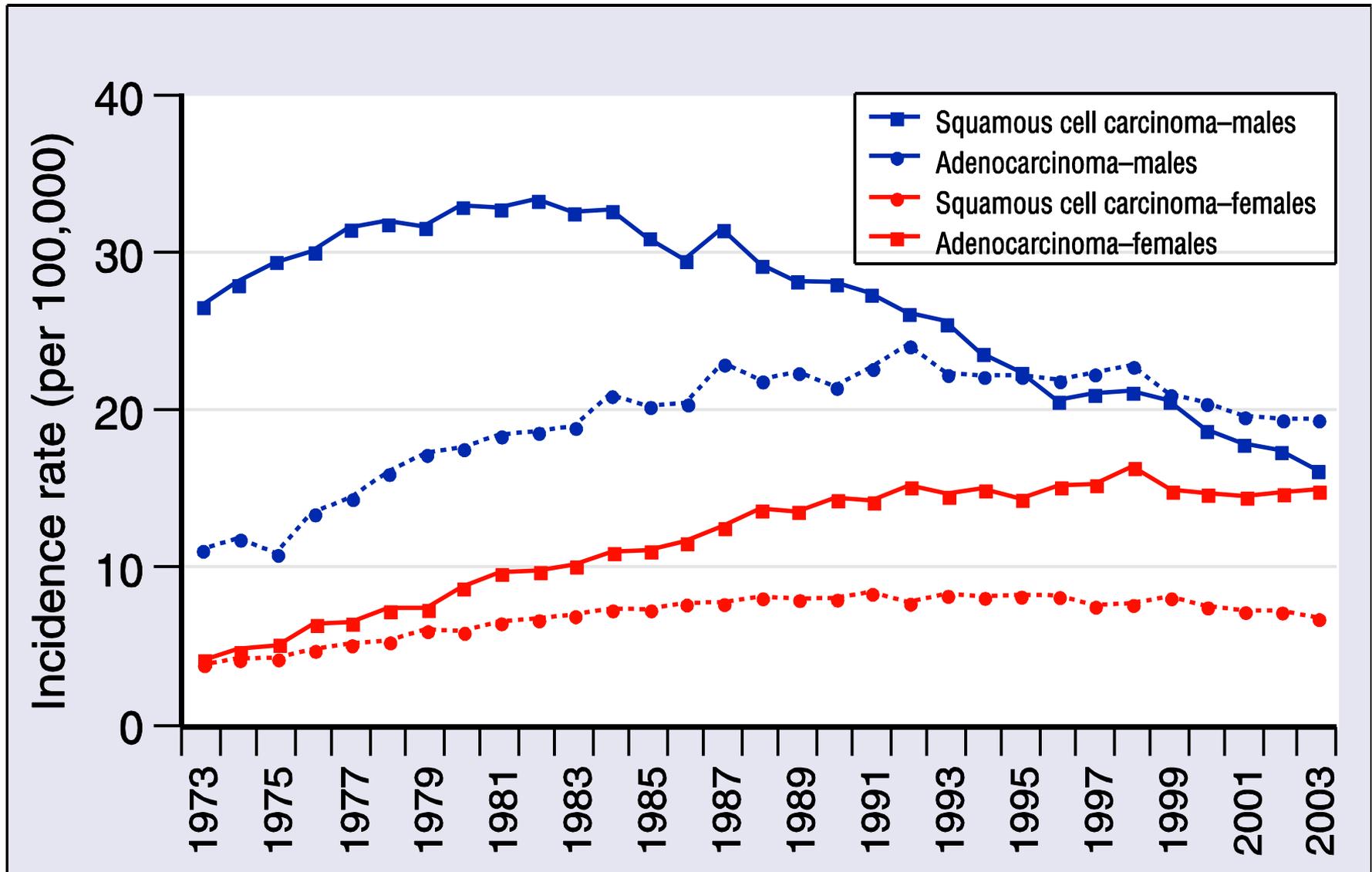


# Annual Deaths Attributed to Tobacco



Center for Disease Control and Prevention, U.S. 2000-2004

# Age-Adjusted NCI SEER Lung Cancer Incidence

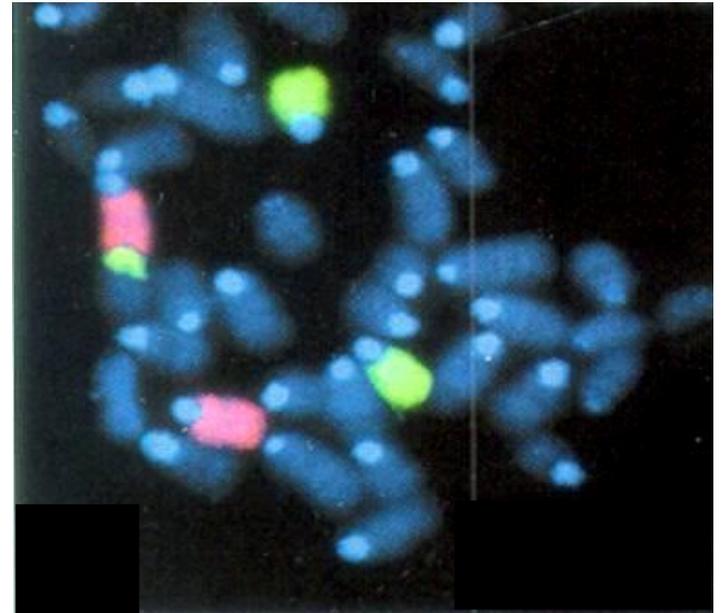


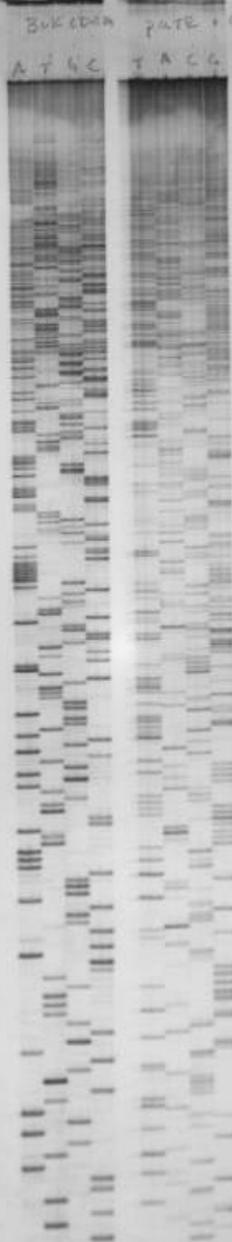
# Cancer is a Disease of the Genome

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It arises from changes  
within the DNA of our cells  
during their lifespan

- Deletions
- Amplifications
- Mutations
- Translocations
- Epigenetic changes
- **Transcriptional/translational regulation**





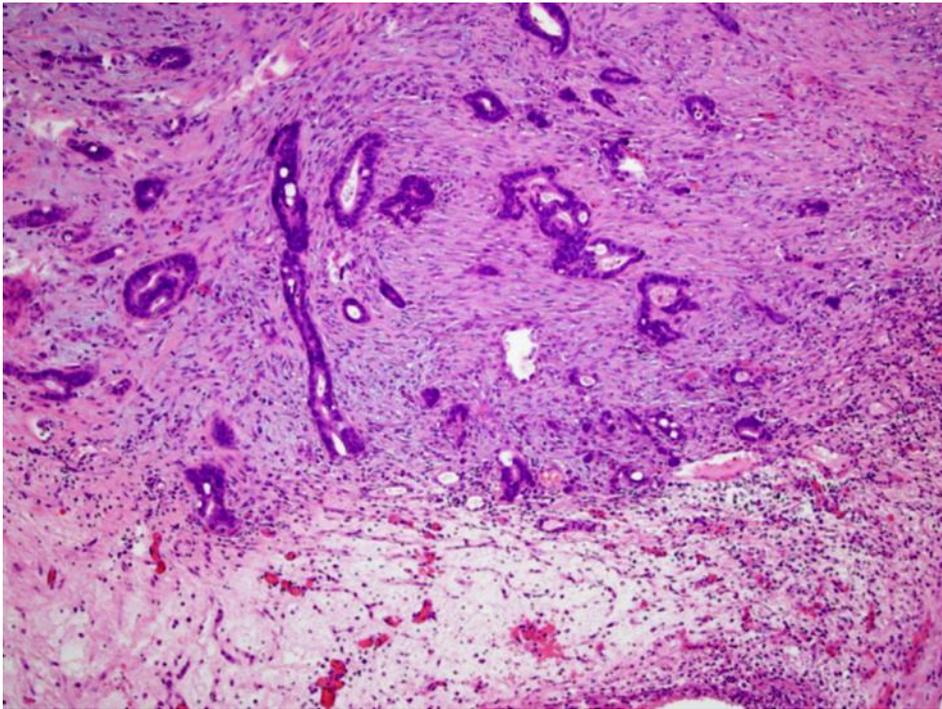
**Mid 1980s**



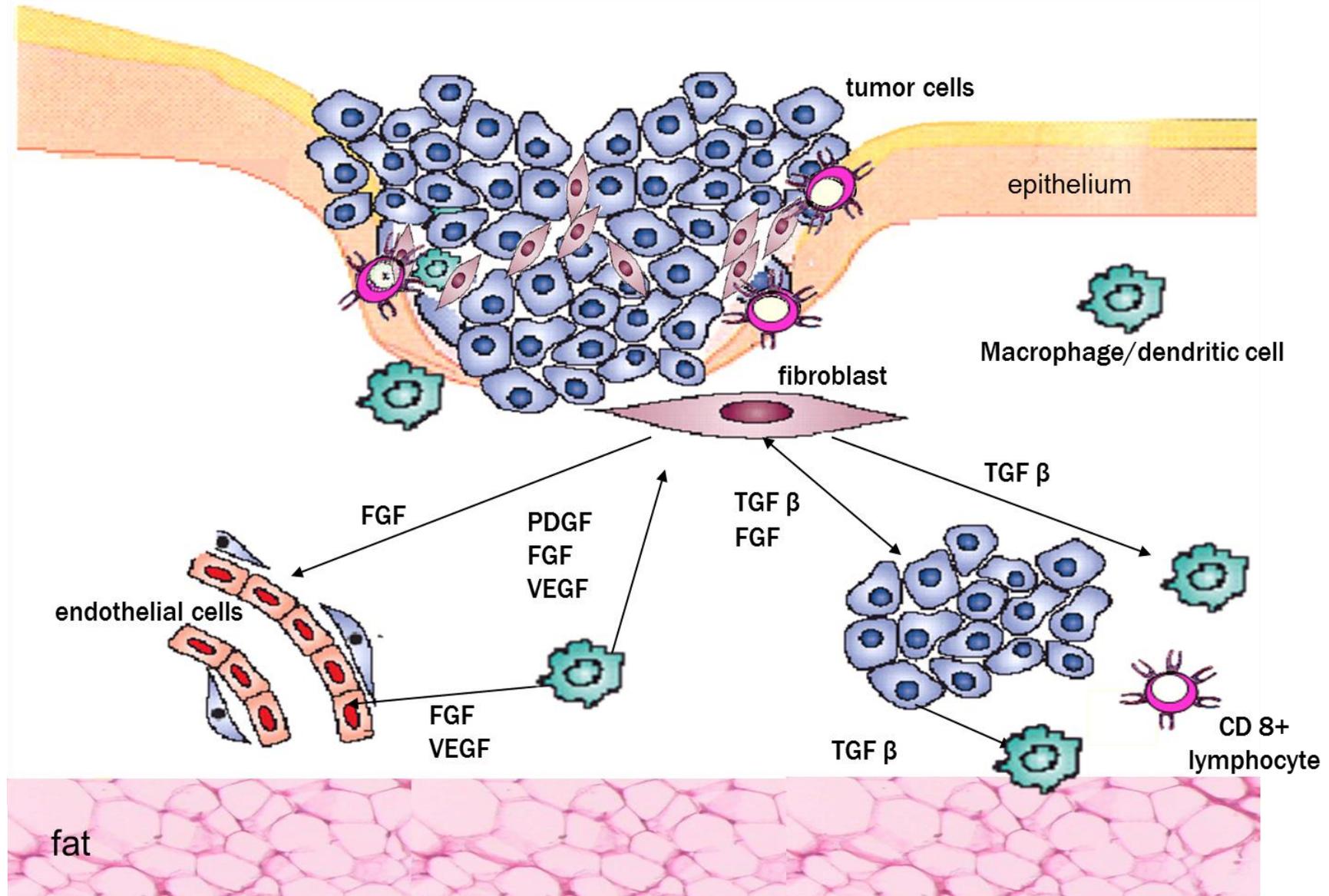
# Tumors as “Organ” Systems

**Tumors are more than just a mass of cancer cells.**

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# Tumors as "Organ" Systems



# Individualized Cancer Care

## Today - some thoughts regarding Genomics and Cancer.

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- **Determining risk**
- **Managing risk**
- **Noval molecular targets**
- **Pharmacogenomics**
- **Each cancer an N of 1**



# Translational Science: The Paradigm Shift

## The 20th Century Paradigm:

Organ site-based, single agent based trials

- Reactive
- Based on gross differences
- Toxic (MTD/DLT)
- Emerging resistance
- Poor life quality

### **Research**

- Human genome
- Genomics
- Proteomics
- Immunology
- Mechanisms
- Rational design

## The New Paradigm:

Multiple, highly targeted agents matched to molecularly selected patients

- Proactive
- Rational/targeted
- Less toxicity
- Biomarker endpoints (subcellular target imaging)
- Significant savings of cost and time

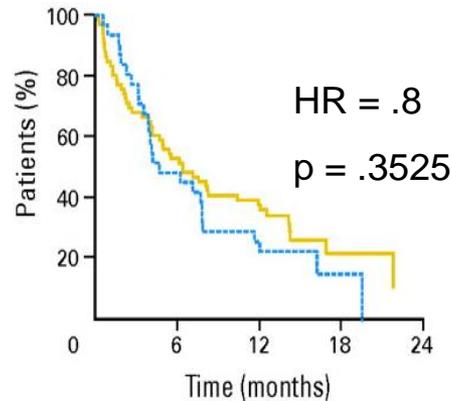
## Creating the repair manual.

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- Derive a functional understanding of the causal defect/dependance; e.g. Wnt, P13K, NF-kB...
  - ✓ Distinguish passenger defects from true drivers.
- Determine dependence of cancer cells and micro-environment cells on genes that are amplified, translocated, mutated or epigenetically altered.
  - ✓ “Oncogene addiction”
- Find genes to which cancer cells are addicted but that are not mutated, translocated or amplified.
  - ✓ “Non-oncogene addiction”

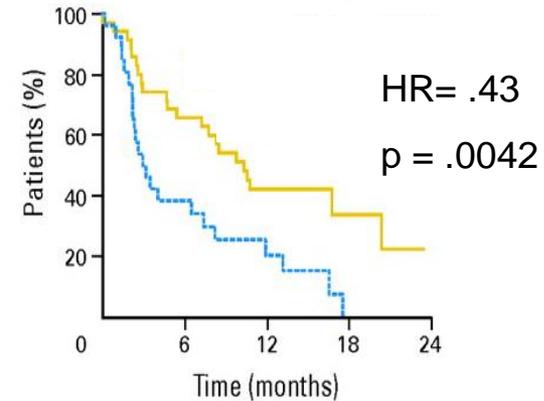
# Kaplan-Meier Survival Curves According to EGFR Copy Number and Impact of Erlotinib

## EGFR Low Copy Number



No. of patients	0	6	12	18	24
Placebo	31	15	8	2	0
Erlotinib	67	35	24	4	1

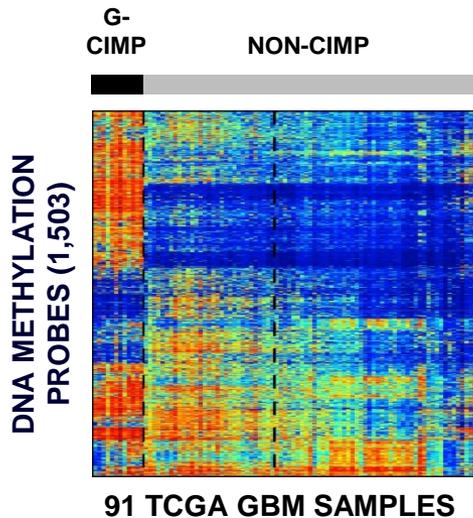
## EGFR High Copy Number



No. of patients	0	6	12	18	24
Placebo	26	9	6	0	0
Erlotinib	35	23	14	4	0

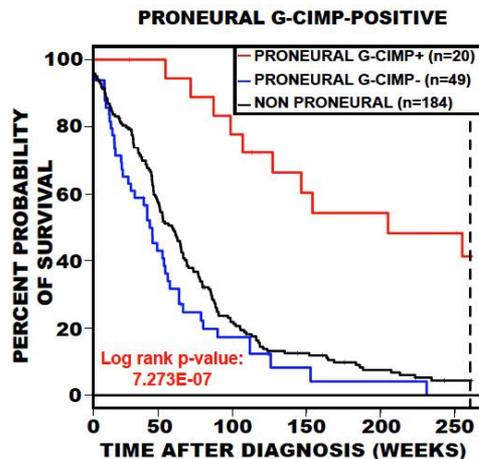
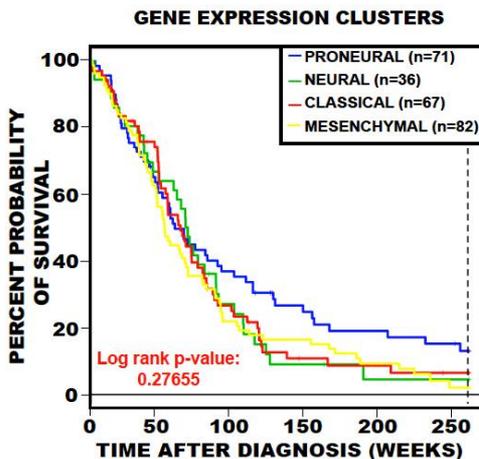


# GBM: CpG Island Methylator Phenotype Linked to *IDH1* Mutation and Better Survival



## G-CIMP.....

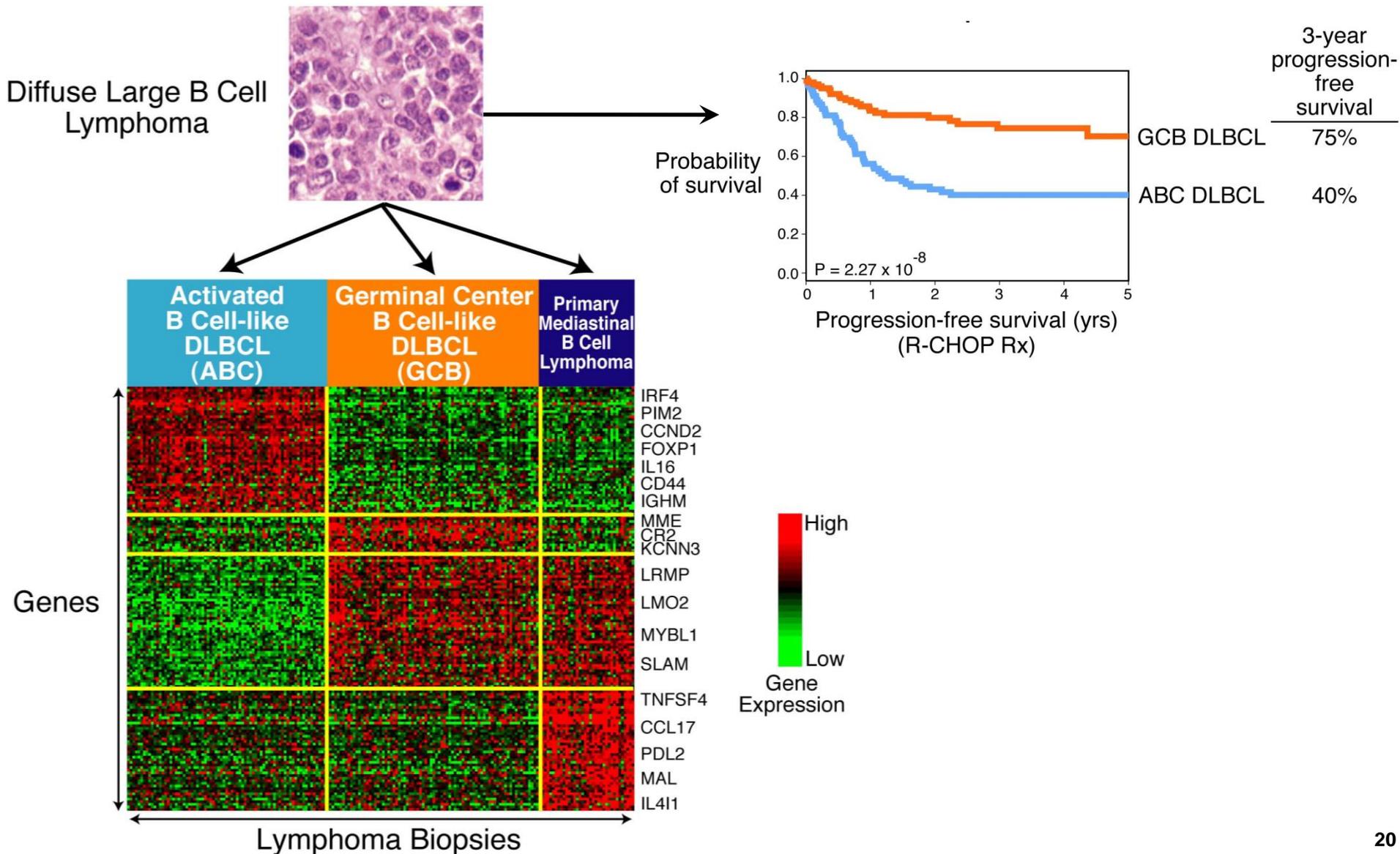
- Occurs in Younger Patients
- Is a Subset of Proneural Expression Subtype
- Is Associated with Better Survival
- Is More Frequent in Low-Grade Gliomas
- Is Not Associated with *MGMT* Methylation
- Is Tightly Linked to *IDH1* Mutation



ALL TUMORS		G-CIMP		TOTAL
		-	+	
IDH1	Wild-type	184	5	189
	Mutant	0	18	18
TOTAL		184	23	207

Noushmehr et al. (2010) Cancer Cell, Online

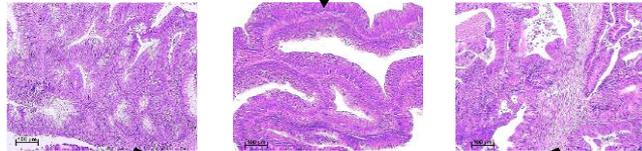
# Dissecting Cancer into Molecularly and Clinically Distinct Subgroups by Gene Expression Profiling



# Individualized Cancer Therapy

## Same Organ Site

**Today**



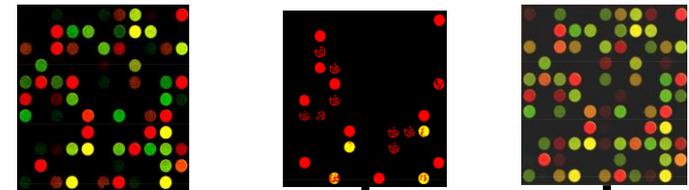
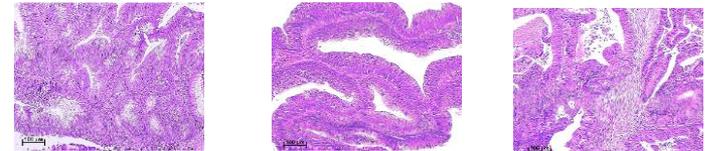
Pt. #1 #2 #3

**Standard Therapy**



**1 in 3 Patients Benefit**

**Future**



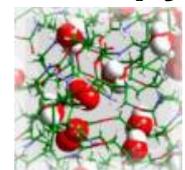
**Therapy 1**



**Therapy 2**



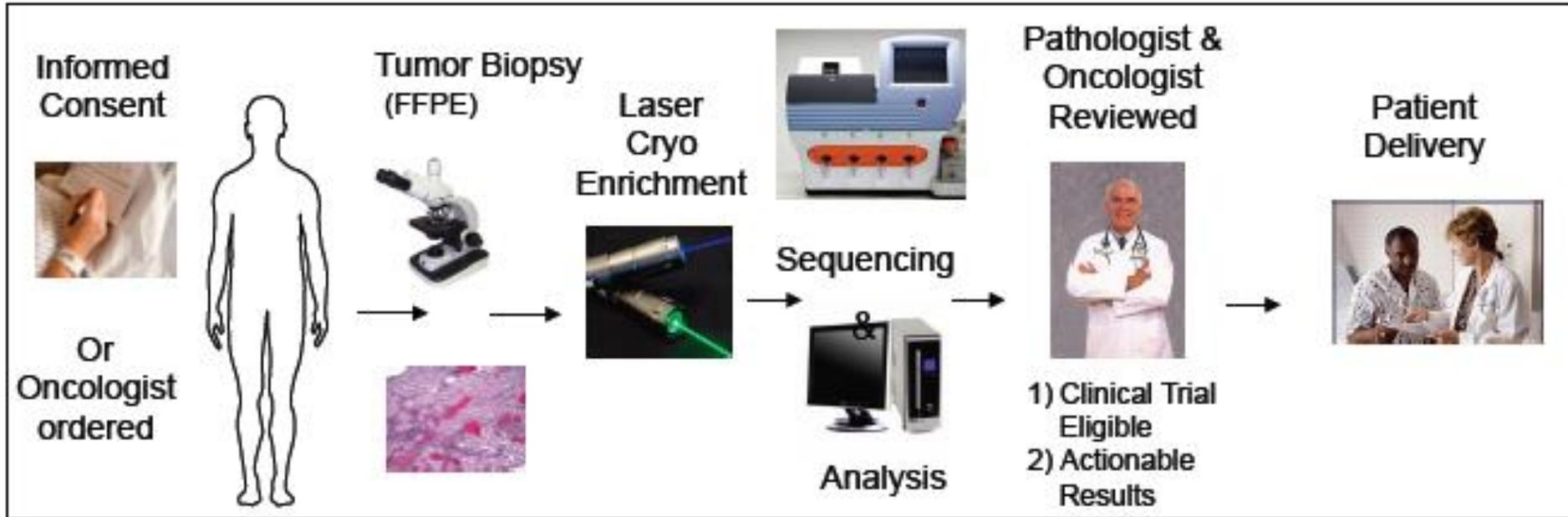
**Therapy 3**



### **Research**

- Genomics
- Proteomics
- Immunology
- Mechanisms
- Rational drug design

# Cancer Diagnosis and Treatment



- **Service offered through three commercial vendors and several academic centers.**
- **Used to diagnose and direct therapy based on genomic identification of biomarkers & targets.**

# ITMI Genomics Data Base

- **Maternal-child genomic studies >2000 families.**
- **WGS @ >40X of mother, father and newborn.**
- **SNP, CNV, SV.**
- **RNAseq expression, CpG methylation.**
- **Clinical study specific data and eHR information.**

## Germline Variations Impacting Cancer Predisposition.

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- **WGS from ancestrally-diverse cohort of 681 healthy adults** – negative pers./fam. Hx.
- **Profiled nonsynonymous variation in 158 genes causally implicated in cancer.**
- **Selected five genes for in depth analysis *BRCA1, BRCA2, KRAS, TP53, and PTEN.***

# Cancer Susceptibility Genes

## Germline Variations Impacting Cancer Predisposition.

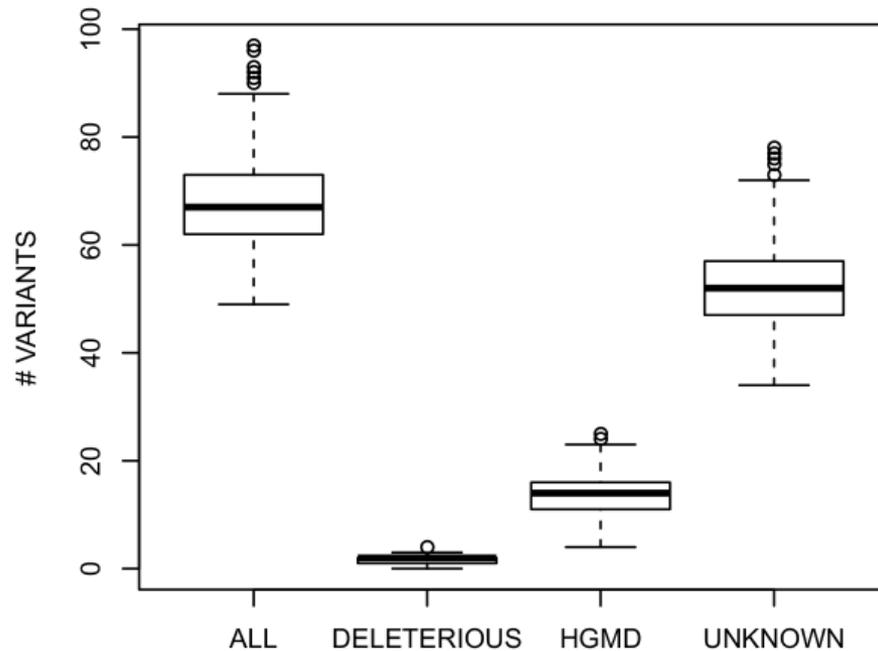
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- **2,688 distinct genetic variants identified within the cohort.**
- **All individuals carry variants that may impact cancer susceptibility.**
- **Average of 68 variants per individual.**
- **Most variants are very rare with 75% found in only one or two individuals.**
- **Allele frequencies vary between ancestral groups.**

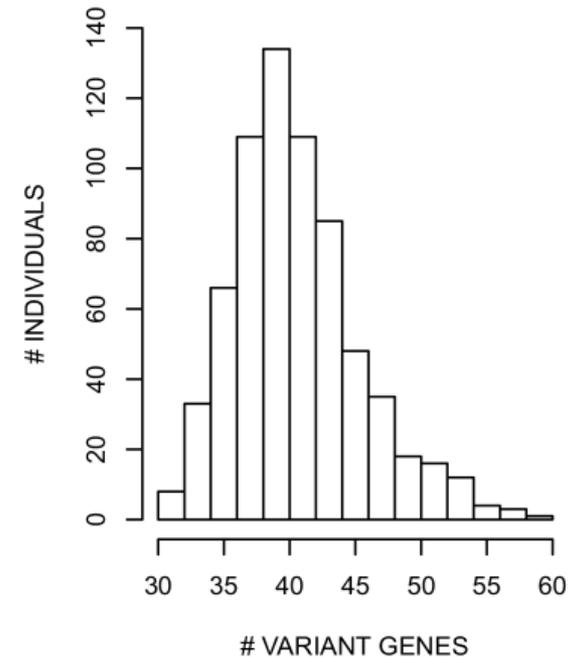
# Cancer Susceptibility Genes

## Germline Variations Impacting Cancer Predisposition.

A



B



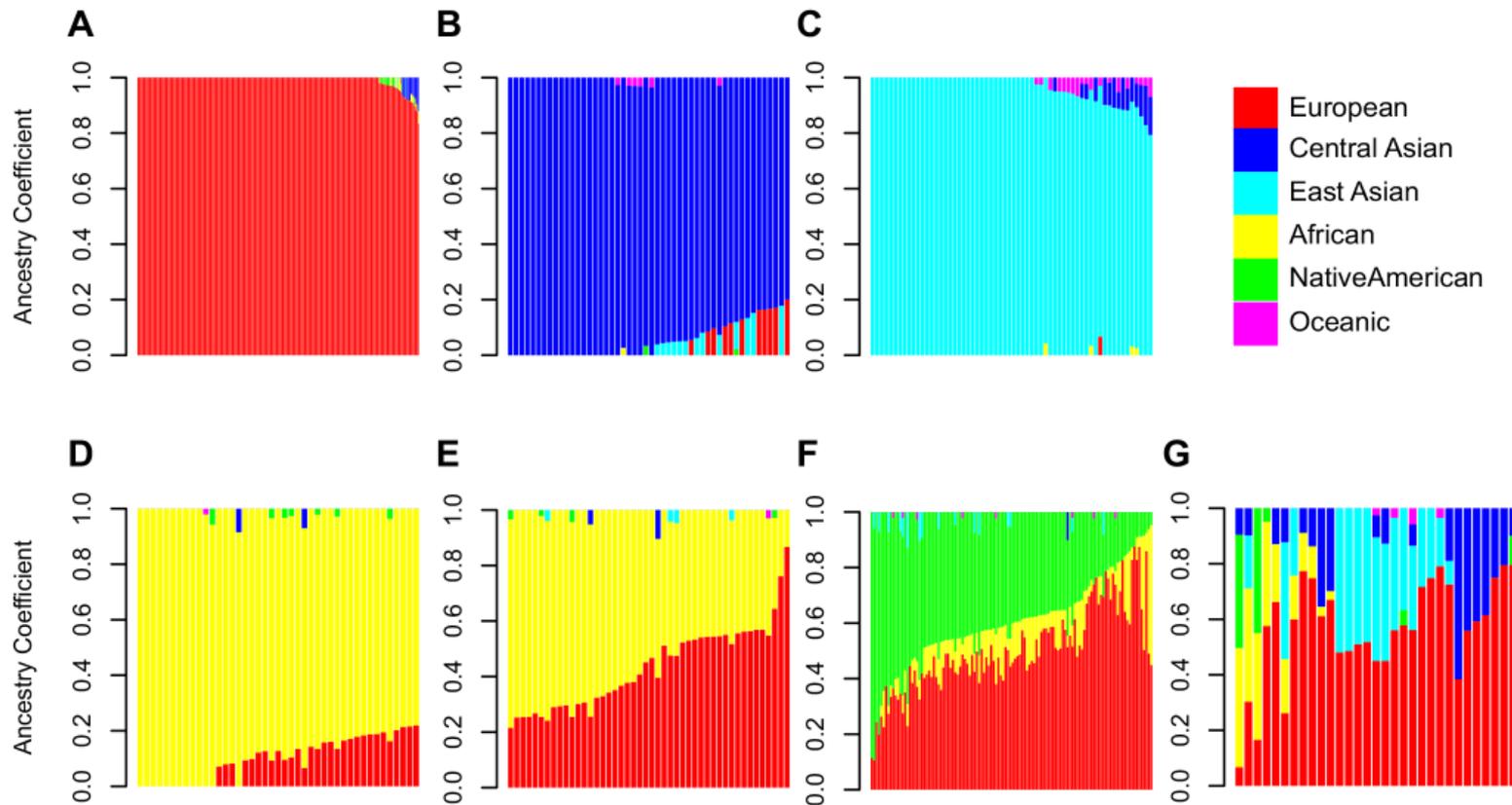
# Cancer Susceptibility Genes

## Ancestry-based subpopulations in study.

<b>Subpopulation</b>	<b># Individuals</b>
African	43
African-European	46
Central Asian	50
East Asian	62
European	331
Hispanic	118
Other	31
	<hr/>
	681

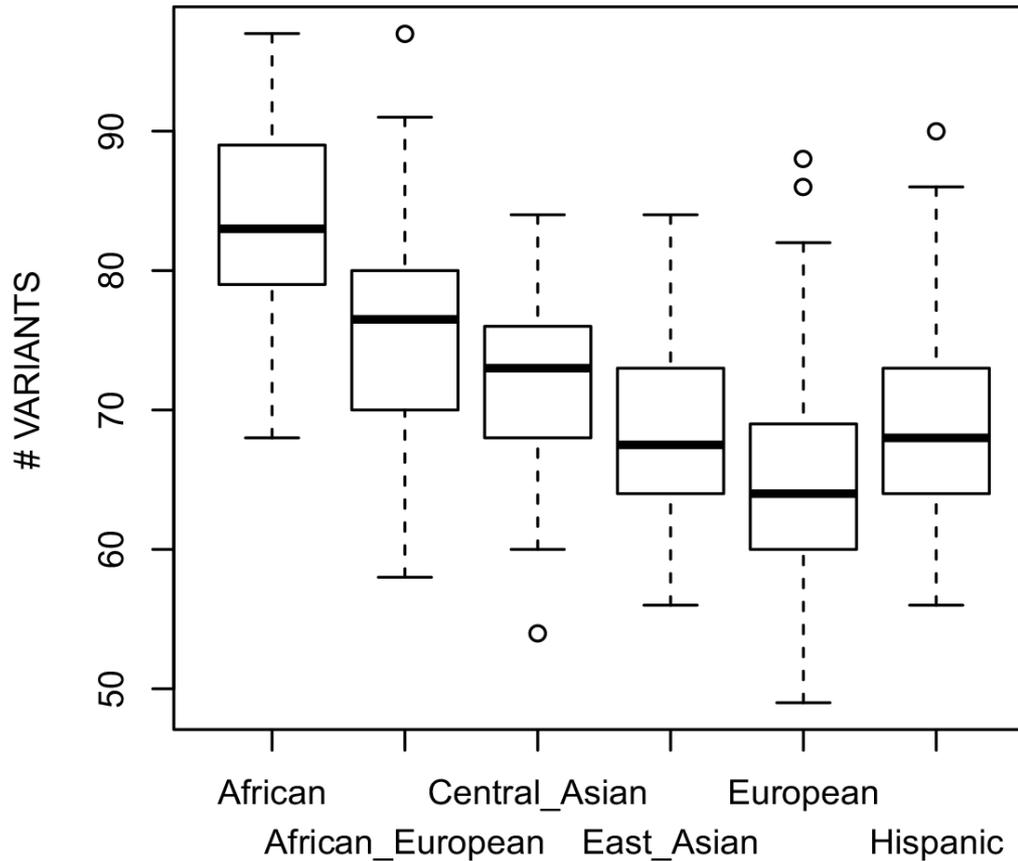
# Cancer Susceptibility Genes

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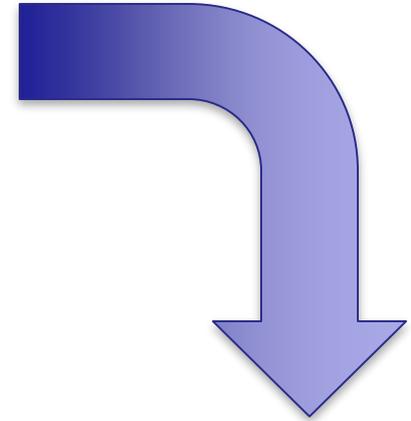
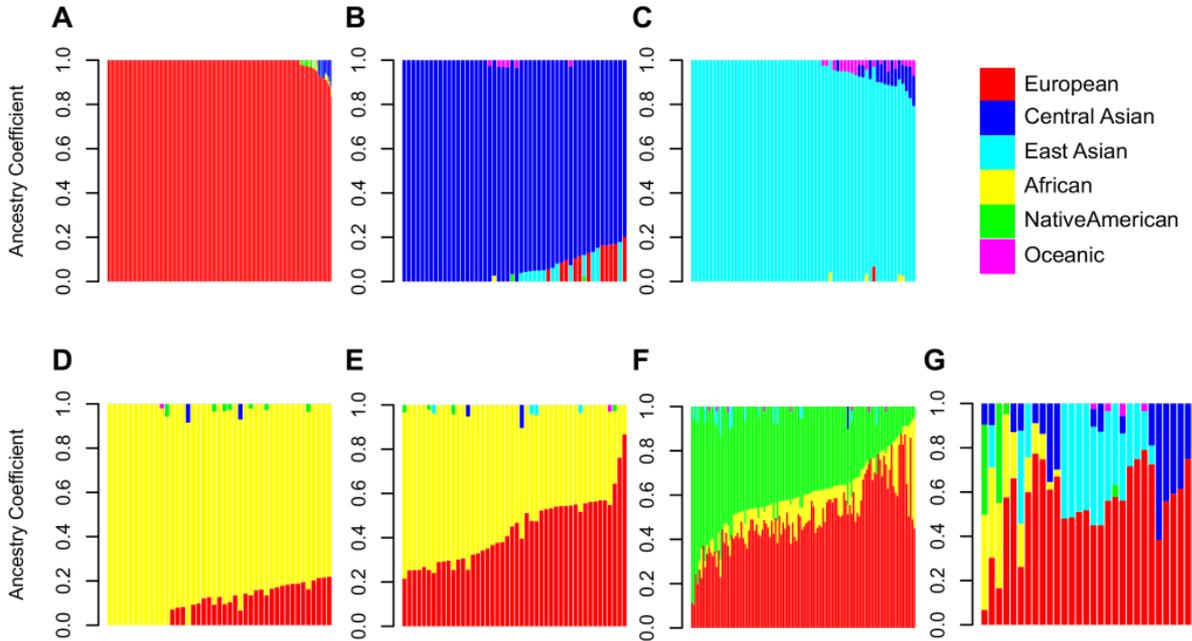


## **Importance of Ancestral Specific Reference Genomes in Genomic Medicine.**

# The Reference Genome

- **The information used in whole genome sequencing to assemble a person's sequence information into a complete genome. (ncbi37)**
- **The reference genome is the basis for identifying relevant variants in a person's genome.**
- **It directly impacts the ability to determine the disease-causing mutations.**
- **The NIH reference genome is not suited to medically relevant sequencing.**

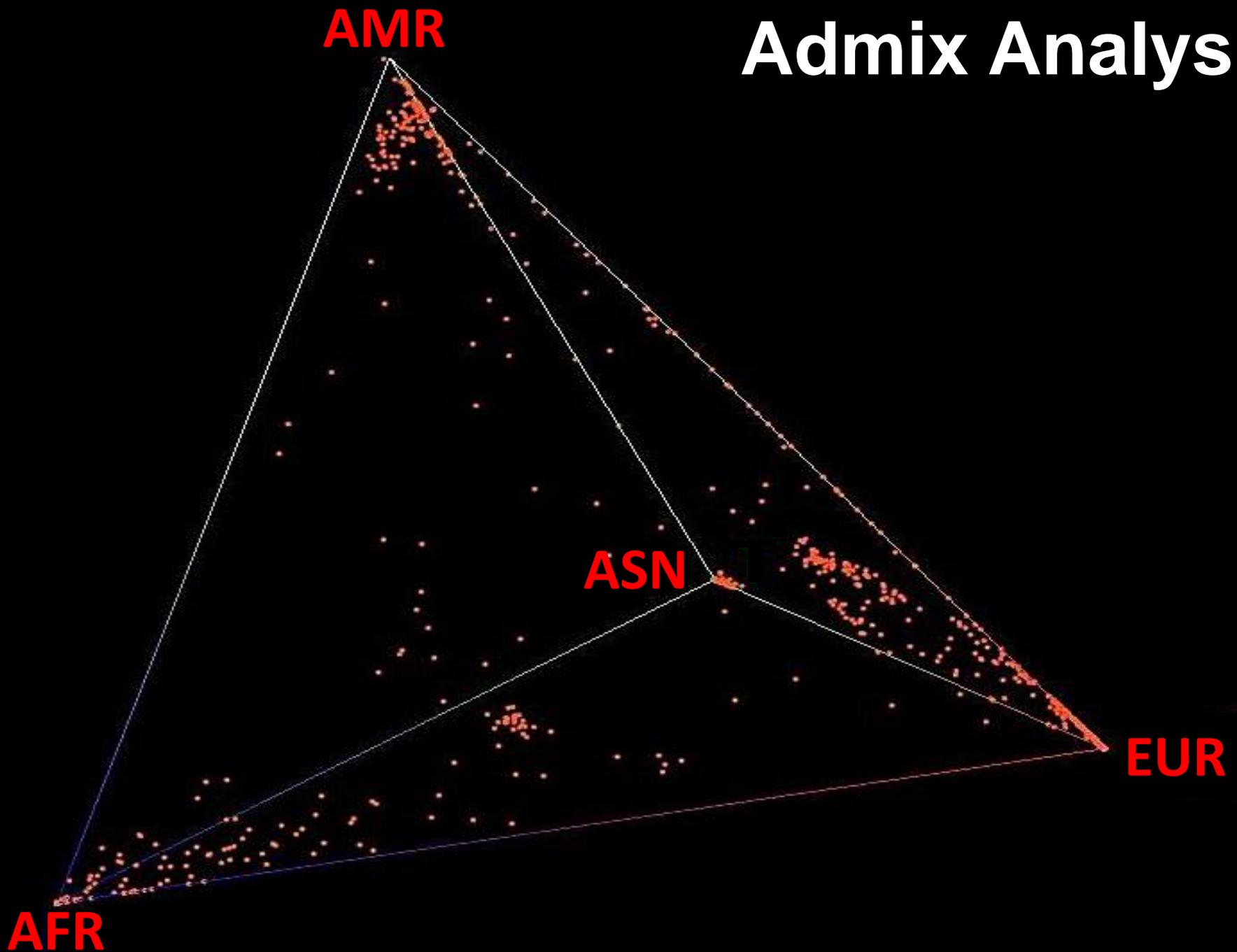
# Admix Analysis



## Build 37 Reference Genome

Donors were recruited by advertisement in *The Buffalo News*, on Sunday, March 23, 1997. DNA was extracted from the blood of 10 male and 10 female volunteers. About 80 percent of the reference genome came from eight people and one male individual accounts for 66 percent of the total reference genome.

# Admix Analysis

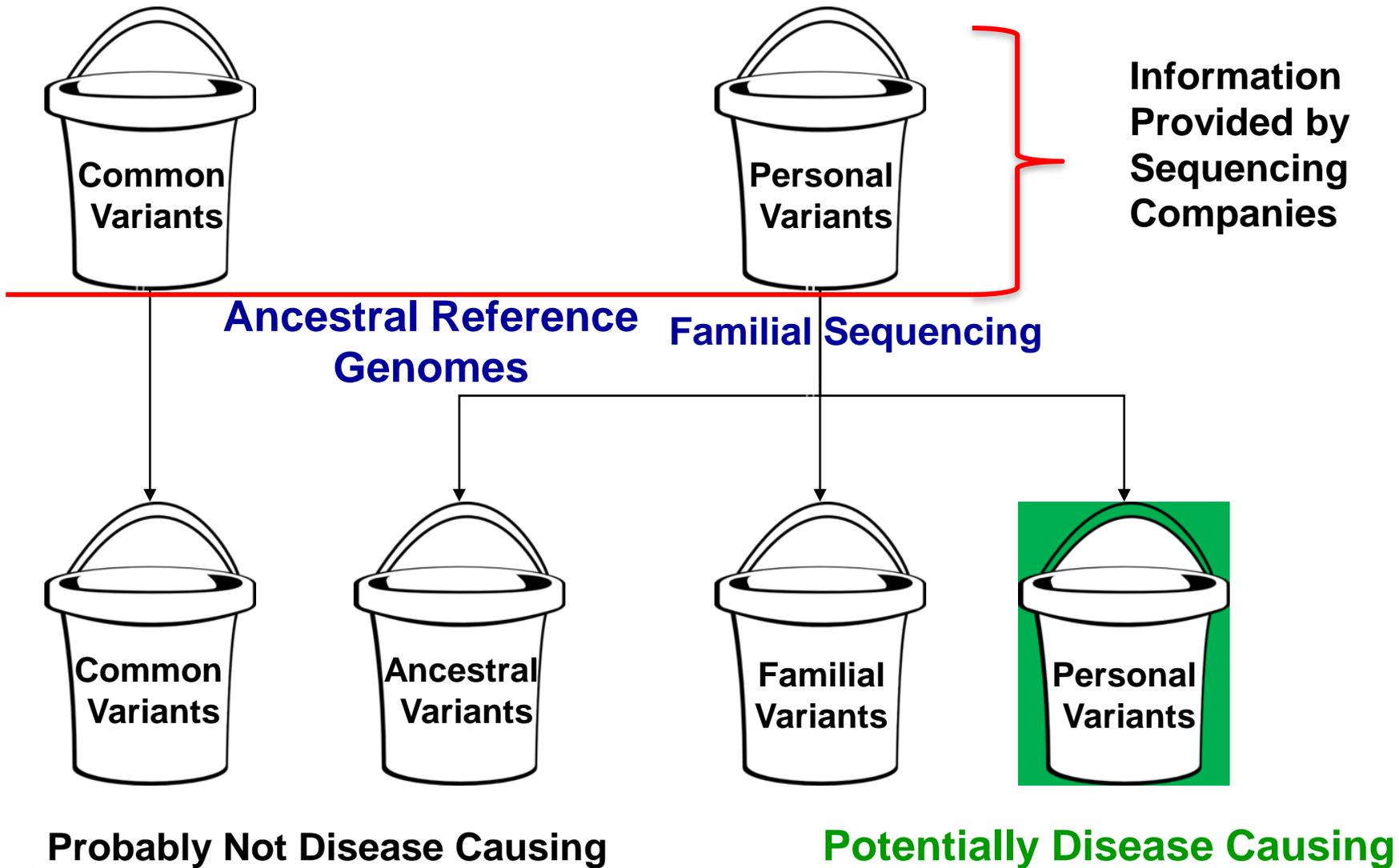


# Pre-Term Birth: Country of Birth



- Country of Birth = 79
- Ancestry currently not used in WGS assembly

# Re-distribution of Personal Variants



# Genomic Medicine Today

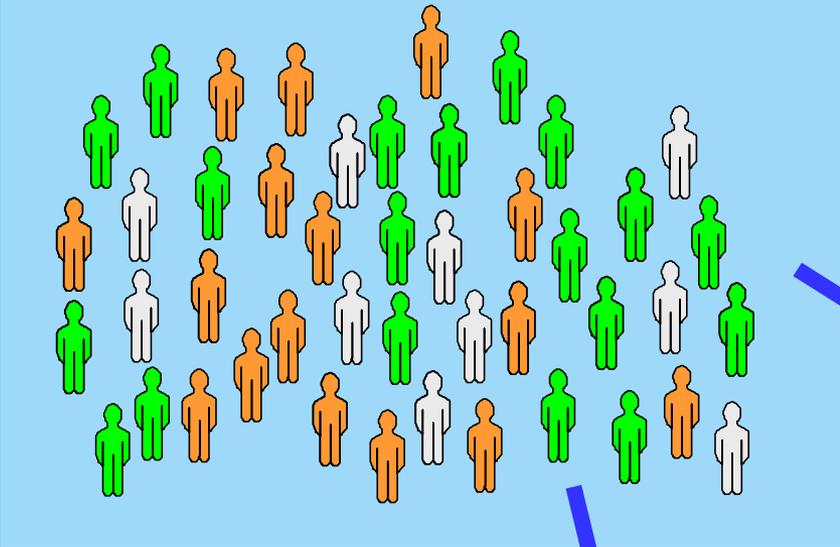
- **Medicine in a reactive discipline, while genomic medicine focuses on prediction (risk) and prevention.**
- **Pharmacogenomics and cancer genomics lead the field in utility.**
- **Through analysis of WGS data we can screen for many actionable genomically based alterations:**
  - **germline cancer predisposition genes, metabolic defects, autism, childhood obesity, cardiovascular disease risk.**
- **Sequence banking will be common practice.**

# Genomic Medicine: Pharmacogenomics

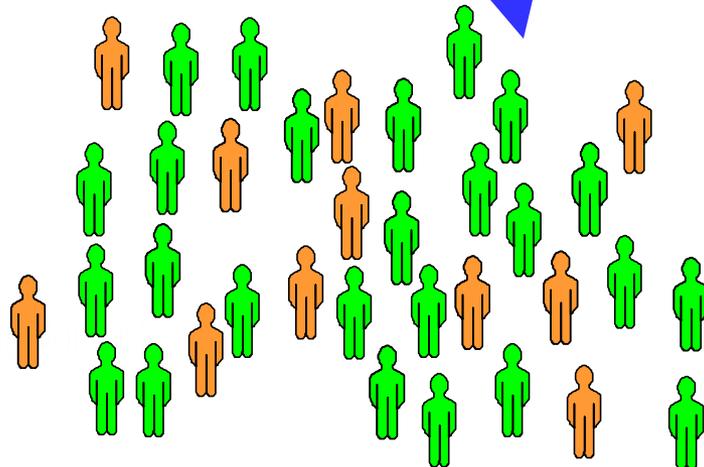
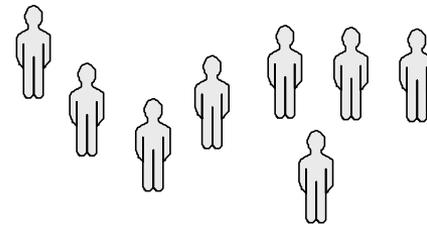
Gene	Total Drugs	Drugs in Key Service Lines	Oncology Drugs	Cardiovascular Drugs	Neuroscicence Drugs
CYP2D6	35	11		Coreg, Toprol, Rhythmol, Innopran Quinidine	Nuedexta, Razadyne Xanaxine, Codine Ultracet
CYP2C19	15	4		Plavix, Effient, Brilinta	Onfi
TPMT	4	4	Platinum Purinethol Thioguanine		Imuran
CYP2C9	3	3	Coumadin	Coumadin	Celebrex
DPD	3	2	Xeloda, Fluorouracil		
G6PD	3	1	Elitek		
UGT1A1	3	2	Camptosar Tasigna		
HLA-B*1502	2	1			Tegretol, Dilantin
NAT1;NAT2	2	1		Bidil	
HLA-B*5701	1	0			

# Pharmacogenomics

All patients with same diagnosis

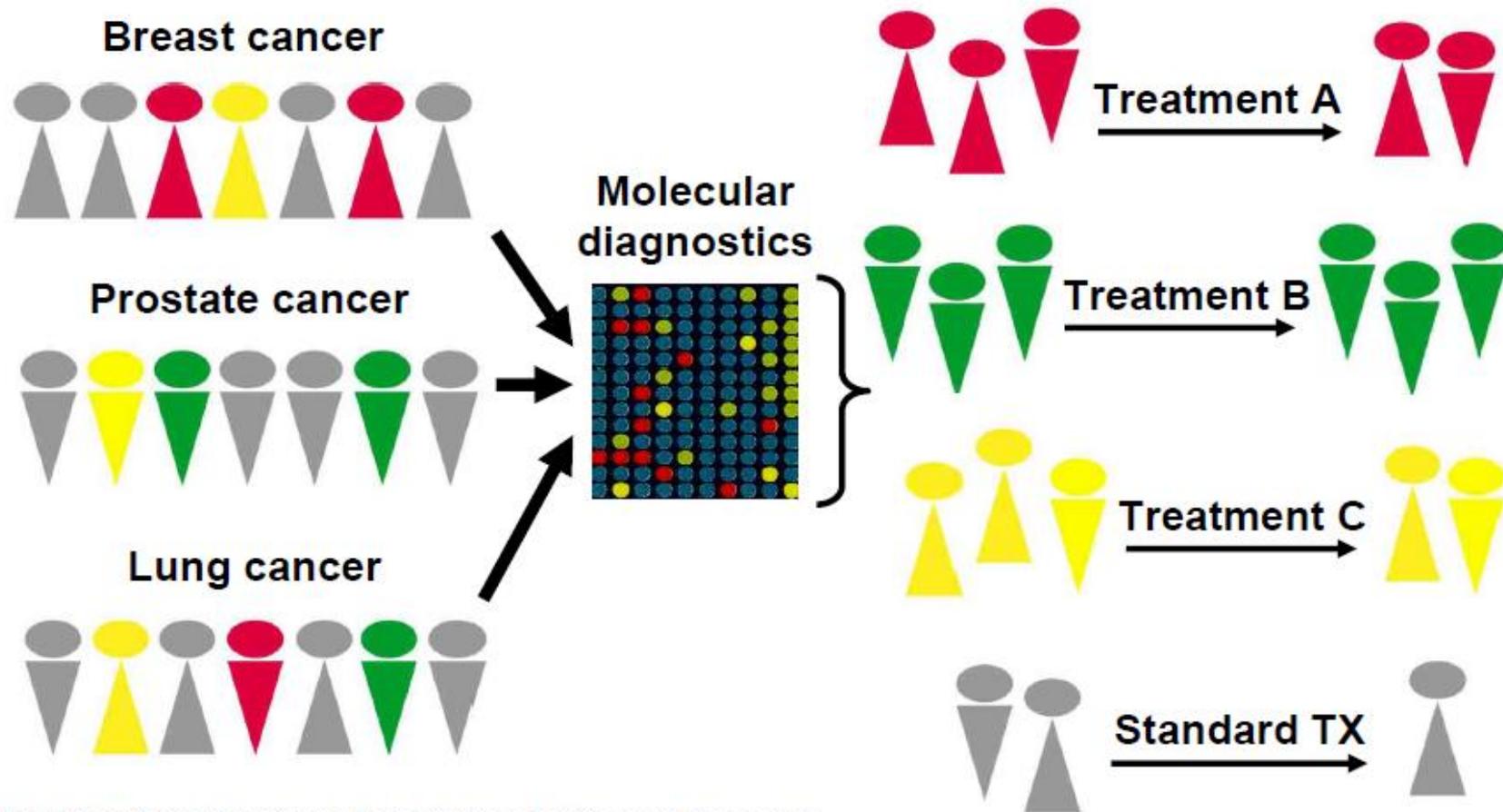


Responders and patients not experiencing severe toxicity



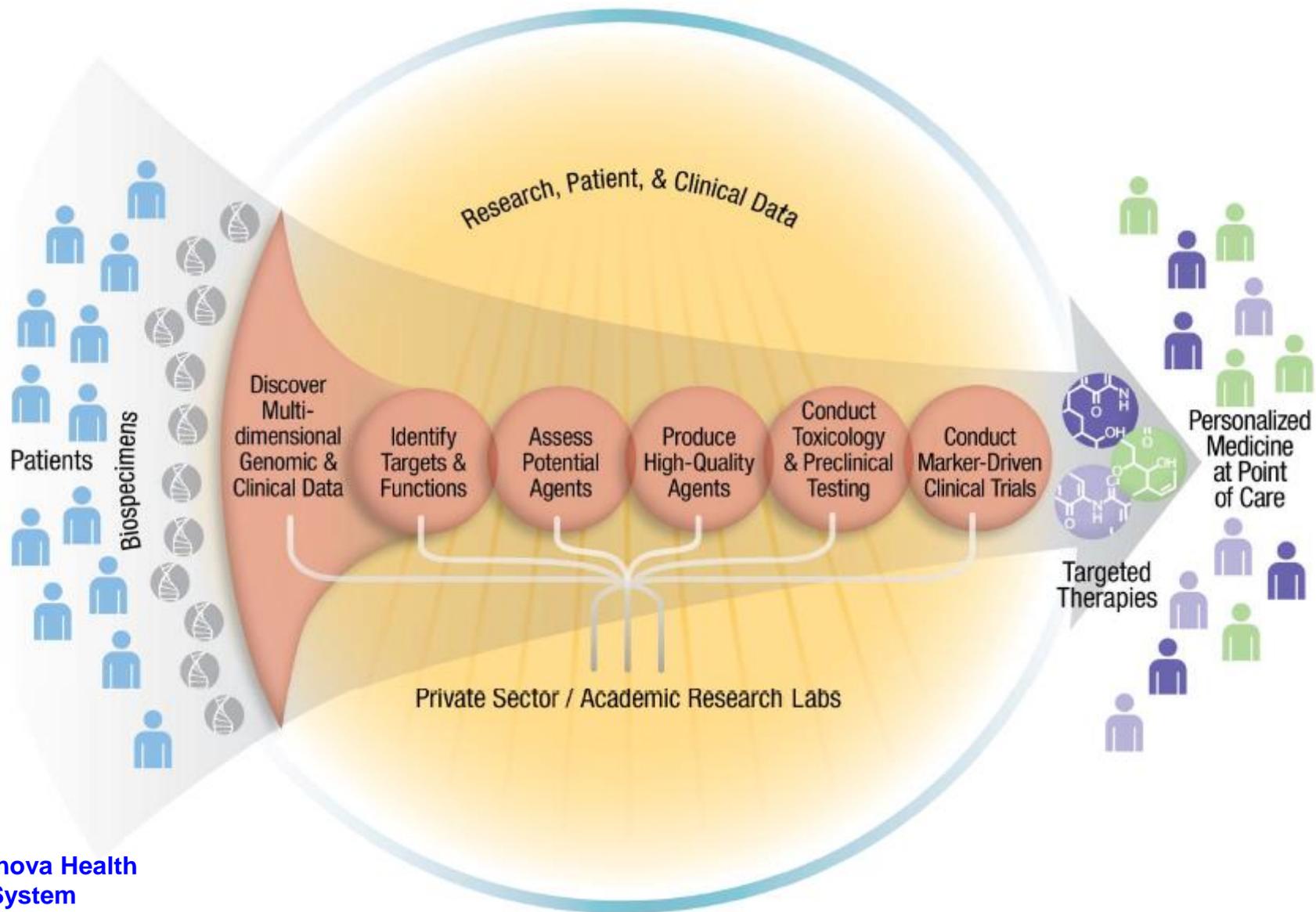
Non-responders  
and toxic responders

# Example: Cancer Therapies



Modified from American Association for Cancer Research

# Genomic Driven Drug Discovery



**The goal of individualized cancer medicine is to use genomic characterization to manage disease risk and to optimize patient therapy.**





